C	nber: 09/787,986A CRF Processing Date: 5/2/200 Edited by: Verified by: Verified by: (STIC s
CI	nanged the margins in cases where the sequence text was "wrapped" down to the next line.
E	lited a format error in the Current Application Data section, specifically:
	lited the Current Application Data section with the actual current number. The number inputted by the plicant was  the prior application data; or other
Αc	Ided the mandatory heading and subheadings for "Current Application Data".
Εc	lited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
Cł	anged the spelling of a mandatory field (the headings or subheadings), specifically:
Cc	rrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
Ins	serted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
	rrected subheading placement. All responses must be on the same line as each subheading. If the plicant placed a response below the subheading, this was moved to its appropriate place.
ln:	serted colons after headings/subheadings. Headings edited included:
De	eleted extra, invalid, headings used by an applicant, specifically:
	eleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of files page numbers throughout text; other invalid text, such as
In	serted mandatory headings, specifically:
	orrected an obvious error in the response, specifically:
E	dited identifiers where upper case is used but lower case is required, or vice versa.
C	orrected an error in the Number of Sequences field, specifically:
Α	"Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
	eted <i>ending</i> stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error to a Patentin bug). Sequences corrected:

\*Examin r: Th abov corr ctions must b communicated to th applicant in th first Office Action. DO NOT send a copy of this form.



DATE: 05/02/2002

TIME: 11:22:43

OIPE

```
Input Set : A:\PTO.AMC.txt
                     Output Set: N:\CRF3\05022002\I787986A.raw
     3 <110> APPLICANT: Lewis, Richard J.
             Alewood, Paul F.
             Sharpe, Iain A.
     7 <120> TITLE OF INVENTION: NOVEL PEPTIDES
     9 <130> FILE REFERENCE: Davies Collison Cave
    11 <140> CURRENT APPLICATION NUMBER: 09/787,986A
C--> 12 <141> CURRENT FILING DATE: 2001-03-23
    14 <160> NUMBER OF SEQ ID NOS: 9
    16 <170> SOFTWARE: PatentIn Ver. 2.1
    18 <210> SEQ ID NO: 1
    19 <211> LENGTH: 13
    20 <212> TYPE: PRT
    21 <213> ORGANISM: Conus marmoreus
    23 <220> FEATURE:
    24 <221> NAME/KEY: PEPTIDE
    25 <222> LOCATION: (12)
    26 <223> OTHER INFORMATION: Xaa at position 12 is 4-hydroxy proline
    28 <400> SEQUENCE: 1
📆-> 29 Asn Gly Val Cys Cys Gly Tyr Lys Leu Cys His Xaa Cys
    33 <210> SEQ ID NO: 2
    34 <211> LENGTH: 13
    35 <212> TYPE: PRT
    36 <213> ORGANISM: Conus marmoreus
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    40 <222> LOCATION: (12)
     41 <223> OTHER INFORMATION: Xaa at position 12 is 4-hydroxy proline
     43 <400> SEQUENCE: 2
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    50 <212> TYPE: PRT
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    56 1
    59 <210> SEQ ID NO: 4
    60 <211> LENGTH: 20
    61 <212> TYPE: DNA
   62 <213> ORGANISM: Artificial Sequence
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/787,986A

- 64 <220> FEATURE:

DATE: 05/02/2002

TIME: 11:22:43 PATENT APPLICATION: US/09/787,986A Input Set : A:\PTO.AMC.txt Output Set: N:\CRF3\05022002\I787986A.raw 65 <223> OTHER INFORMATION: Description of Artificial Sequence: Oligonucleotide probe 68 <220> FEATURE: 69 <221> NAME/KEY: unsure 70 <222> LOCATION: (3) 71 <223> OTHER INFORMATION: n=a/c/g/t 73 <220> FEATURE: 74 <221> NAME/KEY: unsure 75 <222> LOCATION: (6) 76 <223> OTHER INFORMATION: r=a/g 78 <220> FEATURE: 79 <221> NAME/KEY: unsure 80 <222> LOCATION: (9) 81 <223> OTHER INFORMATION: r=a/g 84 <220> FEATURE: 85 <221> NAME/KEY: unsure 86 <222> LOCATION: (12) 87 <223> OTHER INFORMATION: n=a/c/g/t 90 <220> FEATURE: 91 <221> NAME/KEY: unsure 92 <222> LOCATION: (14) 93 <223> OTHER INFORMATION: r=a/g 97 <220> FEATURE: 98 <221> NAME/KEY: unsure 99 <222> LOCATION: (15) 100 <223> OTHER INFORMATION: y=c/t 102 <220> FEATURE: 103 <221> NAME/KEY: unsure 104 <222> LOCATION: (18) 105 <223> OTHER INFORMATION: r=a/g 107 <400> SEQUENCE: 4 20 108 canggrtgrc anaryttrta 111 <210> SEQ ID NO: 5 112 <211> LENGTH: 27 113 <212> TYPE: DNA 114 <213> ORGANISM: Artificial Sequence 116 <220> FEATURE: 117 <223> OTHER INFORMATION: Description of Artificial Sequence: 118 Oligonucleotide probe 120 <400> SEQUENCE: 5 27 121 ccatcctaat acgactcact atagggc 124 <210> SEQ ID NO: 6 125 <211> LENGTH: 23 126 <212> TYPE: DNA 127 <213> ORGANISM: Artificial Sequence 129 <220> FEATURE: 130 <223> OTHER INFORMATION: Description of Artificial Sequence: Oligonucleotide probe 133 <400> SEQUENCE: 6

RAW SEQUENCE LISTING

RAW SEQUENCE LISTING DATE: 05/02/2002 PATENT APPLICATION: US/09/787,986A TIME: 11:22:43

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\05022002\1787986A.raw

23 134 acaggcagaa tgcgctgtct ccc 137 <210> SEQ ID NO: 7 138 <211> LENGTH: 28 140 <212> TYPE: DNA 141 <213> ORGANISM: Artificial Sequence 143 <220> FEATURE: 144 <223> OTHER INFORMATION: Description of Artificial Sequence: Oligonucleotide probe 145 147 <400> SEQUENCE: 7 28 148 aactggaaga attcgcggcc gcaggaat 151 <210> SEQ ID NO: 8 152 <211> LENGTH: 186 153 <212> TYPE: DNA 154 <213> ORGANISM: Conus marmoreus 156 <400> SEQUENCE: 8 157 atgcgctgtc tcccagtctt gatcattctt ctgctgctga ctgcatctgc acctggcgtt 60 158 gttqtcctac cgaagaccga agatgatgtg cccatgtcat ctgtctactg taatggaaag 120 159 agtatcctac gaggaattct gaggaacggt gtgtgctgtg gctataagtt gtgccatcca 180 160 tgttaa 163 <210> SEQ ID NO: 9 164 <211> LENGTH: 61 165 <212> TYPE: PRT 166 <213> ORGANISM: Conus marmoreus 168 <400> SEQUENCE: 9 169 Met Arg Cys Leu Pro Val Leu Ile Ile Leu Leu Leu Thr Ala Ser 170 10 5 1 172 Ala Pro Gly Val Val Leu Pro Lys Thr Glu Asp Asp Val Pro Met 30 20 25 175 Ser Ser Val Tyr Cys Asn Gly Lys Ser Ile Leu Arg Gly Ile Leu Arg 35 40 178 Asn Gly Val Cys Cys Gly Tyr Lys Leu Cys His Pro Cys 179 50 55

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/787,986A

DATE: 05/02/2002 TIME: 11:22:44

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\05022002\I787986A.raw

## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the  $\langle 220 \rangle$  to  $\langle 223 \rangle$  fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 12
Seq#:2; Xaa Pos. 12
Seq#:4; N Pos. 3,12



OIPE

RAW SEQUENCE LISTING

DATE: 04/29/2002

PATENT APPLICATION: US/09/787,986A

TIME: 15:04:59

Input Set : A:\14438.txt

Output Set: N:\CRF3\04292002\I787986A.raw

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3 <110> APPLICANT: Lewis, Richard J.
4 Alewood, Paul F.
5 Sharpe, Iain A.
7 <120> TITLE OF INVENTION: NOVEL PEPTIDES
9 <130> FILE REFERENCE: Davies Collison Cave
11 <140> CURRENT APPLICATION NUMBER: 09/787,986A

C--> 12 <141> CURRENT FILING DATE: 2001-03-23
14 <160> NUMBER OF SEQ ID NOS: 9
16 <170> SOFTWARE: Patentin Ver. 2.1
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## Does Not Comply Corrected Diskette Needed

## **ERRORED SEQUENCES**

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163 <210> SEQ ID NO: 9
    164 <211> LENGTH: 61
    165 <212> TYPE: PRT
    166 <213> ORGANISM: Conus marmoreus
    168 <400> SEQUENCE: 9
    169 Met Arg Cys Leu Pro Val Leu Ile Ile Leu Leu Leu Thr Ala Ser
                                            10
                          5
    172 Ala Pro Gly Val Val Leu Pro Lys Thr Glu Asp Asp Val Pro Met
    173
                    20
                                        25
    175 Ser Ser Val Tyr Cys Asn Gly Lys Ser Ile Leu Arg Gly Ile Leu Arg
          35
                                    40
    178 Asn Gly Val Cys Cys Gly Tyr Lys Leu Cys His Pro Cys
    179
E--> 184(3)
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VERIFICATION SUMMARY

DATE: 04/29/2002

PATENT APPLICATION: US/09/787,986A

TIME: 15:05:00

Input Set : A:\14438.txt

Output Set: N:\CRF3\04292002\1787986A.raw

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:29 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0 L:44 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0 L:108 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0

L:184 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:9